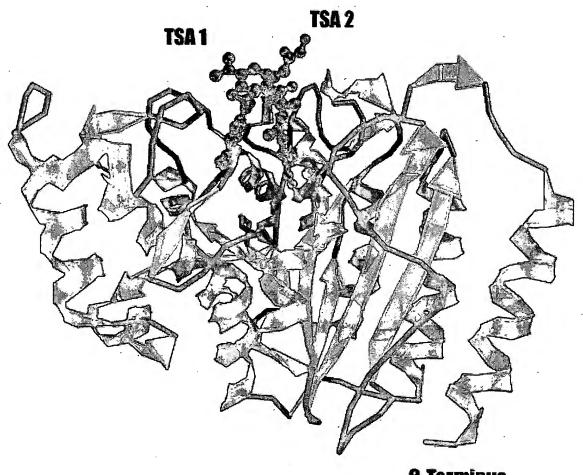
FIGURE 1



6-Terminus

FIGURE 2A

Examples of R₁ and R₂ Substituents

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FIGURE 2A (cont.)

Examples of R_3 and R_4 Substituents

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FIGURE 2A (cont.)

Examples of R₅ Substituents

FIGURE 2B

FIGURE 2C

FIGURE 2C (cont.)

ZZ N R	32 O 35-	ZZ S	R N
\$ // 35-	N Starter Star	N Start	N John St. S
N / St	ZZ S	N-N ZZN R	N-N
R. N-N	7-N 735-	S-N Jos-	R N-N
S-N 12-N	R N / Zz	22 25	The State of the S
N-O Pro-	N-S	R N N	The N
N-N S	N-0 / 5-	N-S	
The North	N-N strains	0-N /	N-N PS-
N S S			

FIGURE 2C (cont.)

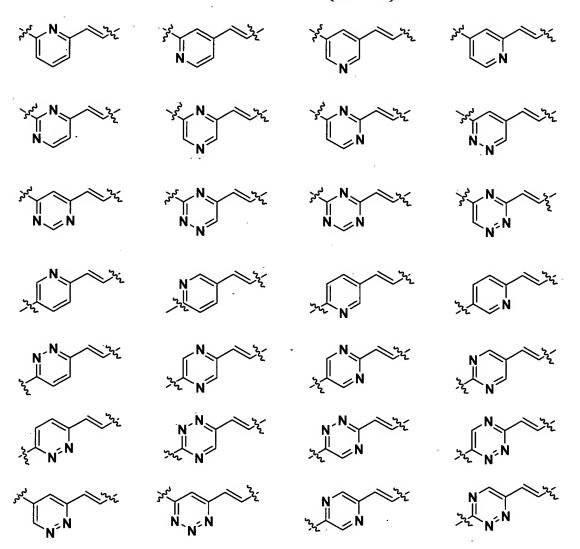


FIGURE 2D

FIGURE 2D (cont.)

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FIGURE 2D (cont.)

FIGURE 3

Amino acid sequence for full length human wild type HDAC8 [SEQ. ID No. 1]

MEEPEEPADSGQSLVPVYIYSPEYVSMCDSLAKIPKRASMVHSLIEAYALHKQMRIVKPK	60
VASMEEMATFHTDAYLQHLQKVSQEGDDDHPDSIEYGLGYDCPATEGIFDYAAAIGGATI	120
TAAQCLIDGMCKVAINWSGGWHHAKKDEASGFCYLNDAVLGILRLRRKFERILYVDLDLH	180
HGDGVEDAFSFTSKVMTVSLHKFSPGFFPGTGDVSDVGLGKGWYYSVNVPIQDGIQDEKY	
YQICESVLKEVYQAFNPKAVVLQLGADTIAGDPMCSFNMTPVGIGKCLKYILQWQLATLI	300
LGGGGYNLANTARCWTYLTGVILGKTLSSEIPDHEFFTAYGPDYVLEITPSCRPDRNEPH	360
RIQQILNYIKGNLKHVV	377

Human cDNA sequence HDAC8 [SEQ. ID No. 2]

ATGGAGGAGCCGGAGAACCGGCGGACAGTGGGCAGTCGCTGGTCCCGGTTTATATCTAT	60
$\tt AGTCCCGAGTATGTCAGTATGTGACTCCCTGGCCAAGATCCCCAAACGGGCCAGTATG$	120
$\tt GTGCATTCTTTGATTGAAGCATATGCACTGCATAAGCAGATGAGGATAGTTAAGCCTAAA$	180
$\tt GTGGCCTCCATGGAGGAGATGGCCACCTTCCACACTGATGCTTATCTGCAGCATCTCCAG$	240
${\tt AAGGTCAGCCAAGAGGGCGATGATGATCATCCGGACTCCATAGAATATGGGCTAGGTTAT}$	300
${\tt GACTGCCCAGCCACTGAAGGGATATTTGACTATGCAGCAGCTATAGGAGGGGCTACGATC}$	360
${\tt ACAGCTGCCCAATGCCTGATTGACGGAATGTGCAAAGTAGCAATTAACTGGTCTGGAGGG}$	420
${\tt TGGCATCATGCAAAGAAGAAGAAGCATCTGGTTTTTGTTATCTCAATGATGCTGTCCTG}$	480
${\tt GGAATATTACGATTGCGACGGAAATTTGAGCGTATTCTCTACGTGGATTTGGATCTGCAC}$	540
${\tt CATGGAGATGGTGTAGAAGACGCATTCAGTTTCACCTCCAAAGTCATGACCGTGTCCCTG}$	600
${\tt CACAAATTCTCCCAGGATTTTTCCCAGGAACAGGTGACGTGTCTGATGTTGGCCTAGGG}$	660
${\tt AAGGGATGGTACTACAGTGTAAATGTGCCCATTCAGGATGGCATACAAGATGAAAAATAT}$	720
${\tt TACCAGATCTGTGAAAGCGTACTAAAGGAAGTATACCAAGCCTTTAATCCCAAAGCAGTG}$	780
$\tt GTCTTACAGCTGGGAGCTGACACAATAGCTGGGGATCCCATGTGCTCCTTTAACATGACT$	840
CCAGTGGGAATTGGCAAGTGTCTTAAGTACATCCTTCAATGGCAGTTGGCAACACTCATT	900
$\tt TTGGGAGGAGGCTATAACCTTGCCAACACGGCTCGATGCTGGACATACTTGACCGGG$	960
$\tt GTCATCCTAGGGAAAACACTATCCTCTGAGATCCCAGATCATGAGTTTTTCACAGCATAT$	1020
${\tt GGTCCTGATTATGTGCTGGAAATCACGCCAAGCTGCCGGCCAGACCGCAATGAGCCCCAC}$	1080
CGAATCCAACAAATCCTCAACTACATCAAAGGGAATCTGAAGCATGTGGTCTAG	1134

FIGURE 4

Amino acid sequence for residues 1-377 of HDAC8 with a cleavable N-terminal 6x-histidine tag [SEQ. ID No. 3]

(6x-histidine tag underlined)

<u>MHHHHHHPMEEPEEPADSGQSLVPVYIYSPEYVSMCDSLAKIPKRASMVHSLIEAYALHK</u>	60
${\tt QMRIVKPKVASMEEMAAFHTDAYLQHLQKVSQEGDDDHPDSIEYGLGYDCPATEGIFDYA}$	120
AAIGGATITAAQCLIDGMCKVAINWSGGWHHAKKDEASGFCYLNDAVLGILRLRRKFERI	180
$\verb LYVDLDLHHGDGVEDAFSFTSKVMTVSLHKFSPGFFPGTGDVSDVGLGKGRYYSVNVPIQ $	240
DGIQDEKYYQICESVLKEVYQAFNPKAVVLQLGADTIAGDPMCSFNMTPVGIGKCLKYIL	300
${\tt QWQLATLILGGGGYNLANTARCWTYLTGVILGKTLSSEIPDHEFFTAYGPDYVLEITPSC}$	360
RPDRNEPHRIQQILNYIKGNLKHVV	385

FIGURE 5